

Table S1. Data collection and refinement statistics for the LIN-41 filamin domain

<i>Data collection</i>	
Space group	C222 ₁
Cell const. <i>a</i> , <i>b</i> , <i>c</i> [Å]	44.78, 52.11, 101.31
Wavelength [Å]	1.000
Resolution range [Å] ^a	50.0-1.68 (1.72-1.68)
Unique reflections	13883
Completeness [%] ^a	99.9 (100.0)
Multiplicity	7.0
<i>R</i> _{sym} [%] ^a	4.5 (87.8)
<i>I</i> / σ (<i>I</i>) ^a	20.8 (1.9)
CC(1/2) [%] ^a	99.9 (69.3)
<i>Refinement</i>	
Resolution range [Å]	50.0-1.68
Reflections (all)	13854
Reflections (test set)	693 (5.0 %)
<i>R</i> _{crys} [%]	21.9
<i>R</i> _{free} [%]	24.4
RMSDs	
Bond lengths [Å]	0.01
Bond angles [°]	1.11
Ramachandran plot	
[%]	
Allowed	100
Outliers	0

^aValues in parentheses refer to the highest resolution shell